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DESCRIPTION

Luciferase and Methods for Measuring Intracellular ATP using the same

Technical Field

The present invention relates to novel luciferase having resistance to a surfactant and a method for measuring intracellular ATP using the same.

Background Art

Intracellular ATP is routinely measured for determining the presence of cells in a sample or the number of cells in the fields of food sanitation, biology, clinical examinations, medical science, ultrapure water, and environmental science. A general method for measuring intracellular ATP comprises the steps of adding an ATP extraction reagent containing a surfactant as an effective component to a sample containing cells, extracting intracellular ATP, adding a luminescence reagent containing luciferase into the sample, and then measuring the total amount of light emitted.

Luciferase is an enzyme that catalyzes luminescence reaction of luciferin, which is a substrate, in the presence of ATP and magnesium ion. Luciferase used in a method for measuring intracellular ATP includes those derived from firefly species, such as GENJI firefly (Luciola cruciata), HEIKE firefly (Luciola lateralis), North American firefly and Russian firefly, etc.

Intracellular ATP can be extracted by adding an ATP extraction reagent to a sample containing cells and then stirring the sample.

To make full use of the capabilities of the extraction reagent, preferably the reaction agent is added so that the concentration of a surfactant becomes 0.05% or more of the mixture of the sample and the extraction reagent. However, a condition where the concentration of the surfactant is 0.05% or more, this inhibits significantly the enzyme

reaction in the process of measuring ATP concentration by bioluminescence. Thus the sensitivity and accuracy of measurement are largely impaired. This is because a surfactant at such a high concentration lowers luciferase activity.

For example, North American firefly luciferase activity decreases to about 20% in the presence of 0.1% benzalkonium chloride (See Table 1).

On the other hand, inhibition of the bioluminescent reaction can be reduced with a lower concentration of surfactant. However, in this case the extraction efficiency for ATP would be insufficient.

A method wherein cyclodextrin or its derivative is used is a known method for suppressing the inhibition of luminescence reaction by a surfactant (Japanese Patent Application Laid-Open No. 6-504200).

Among methods for measuring intracellular ATP wherein intracellular ATP is extracted by allowing a sample to contact with a surfactant and subsequently ATP is measured by luciferin-luciferase bioluminescent reaction method, a method for measuring intracellular ATP characterized by the application of the bioluminescent reaction method after allowing a sample, from which ATP is extracted, to contact with cyclodextrin (Japanese Patent Application Laid-Open Publication No. 7-203995) is also known.

There has been no attempt so far to suppress the inhibition of bioluminescent reaction due to a surfactant focusing on luciferase.

The purpose of the invention is to provide a novel luciferase having anti-surfactant resistance, whose activity is not impaired by the presence of a surfactant at a high concentration. The other purpose of the invention is to provide a method, comprising the steps of extracting intracellular ATP using a surfactant and measuring intracellular ATP by bioluminescent reaction using a luciferase, which can lower the inhibition of

bioluminescent reaction due to a surfactant without a decrease in efficiency in extracting intracellular ATP.

In the context of this Specification, the term "suppress" is used to describe significant reduction of the inhibition of the luminescence reaction by a surfactant and the complete elimination of this inhibition.

Disclosure of the Invention

The present invention relates to a luciferase having anti-surfactant resistance.

The luciferase having resistance to a surfactant includes a luciferase, wherein an amino acid at the 490-position, or an amino acid corresponding to the amino acid at 490-position of GENJI firefly or HEIKE firefly is substituted by an amino acid other than glutamic acid, e.g., lysine, in the amino acid sequence of a wild-type firefly luciferase.

Further, the luciferase having resistance to a surfactant includes a polypeptide consisting of (a) or (b):

- (a) A polypeptide consisting of the amino acid sequence shown in SEQ ID NO:4,
- (b) A polypeptide comprising additions, deletions, or substitutions of one or more of amino acids in the polypeptide of (a), and having luciferase activity resistant to a surfactant, or
- a polypeptide consisting of (a) or (b):
- (a) A protein consisting of an amino acid sequence shown in SEQ ID NO:6,
- (b) A protein comprising additions, deletions, or substitutions of one or more of amino acids in the polypeptide of (a), and having luciferase activity resistant to a surfactant.

Further, the present invention relates to a luceferase gene encoding the luciferase having resistance to a surfactant.

Furthermore, the present invention relates to a recombinant vector containing the luciferase gene encoding the luciferase having resistance to a surfactant.

The present invention also relates to a transformant containing the recombinant vector.

In addition, the present invention relates to a method for producing the luciferase, comprising the steps of culturing the recombinant in a medium, and collecting luciferase with resistance to a surfactant from the culture product.

Moreover the present invention relates to a method for measuring intracellular ATP, comprising the steps of a first step wherein ATP is extracted in the presence of a surfactant from cells in a sample; a second step wherein a luminescence reagent containing luciferase is added to the extracted ATP solution so as to cause light emission; and a third step wherein the light emission is measured, and characterized in that luciferase having resistance to a surfactant is used.

This specification encompasses the description and/or drawings given in Japanese Patent Application No. H09-361022.

Brief Description of Drawings

Figure 1 shows a production processes for a mutant luciferase HIK.

Figure 2 shows change with time of light emission from natural type luciferase.

Figure 3 shows a comparative resistance against benzalkonium chloride of mutant luciferase.

Figure 4 shows a comparative resistance against benzetonium chloride of mutant luciferase.

Detailed Description of the Invention

The present invention will now be described in detail.

[Luciferase having resistance to surfactant]

Luciferase having resistance to a surfactant according to the present invention is as described below.

The term "having resistance to a surfactant" corresponds to any one of the following features.

- (1) When compared to known luciferase, the luciferase of the present invention leads to an increased initial amount of light emitted in the presence of a surfactant. Here the term "compare" means, for example, where the luciferase of the present invention is produced by introducing mutation into an amino acid sequence of known luciferase, to compare light emission from luciferase before and after the introduction of a mutation.
- (2) When compared to known luciferase, the luciferase of the present invention shows a gentle decrease in its activity in the presence of a surfactant.
- (3) The luciferase of the present invention has the remaining activity of more than 85% in the present of 0.4% surfactant.

Hereinafter "luciferase having resistance to a surfactant" is referred to as "surfactant – resistant luciferase."

The term "activity" means the catalytic activity of bioluminescent reaction. Further any surfactant can be used in the present invention so far as it can be used in the measurement system for intracellular ATP. These surfactants include an anionic surfactant, cationic surfactant, ampholytic surfactant, non-ionic surfactant. A specific reagent is benzalkonium chloride or benzetonium chloride containing quaternary ammonium salt as a major component.

The luciferase of the present invention can be prepared from luminescence organs of luminescent organisms. The luminescent organisms include luminescent insects and

luminescent bacteria. The luminescent insects include those belonging to the order Cleoptera, such as those belonging to the family firefly and the family Pyrophorus. Specific examples include GENJI firefly, HEIKE firefly, North American firefly, Russian firefly, Pynophorus plagiophthalamus, Arachnocampa luminosa, and Rail worm. Further the luciferase of the present invention is obtained by cloning a luciferase gene from the luminescent organism and allowing the gene to express using an appropriate vector – host system.

Moreover, the luciferase of the present invention can be obtained by introducing mutation such as additions, deletions, and substitutions into an amino acid sequence of well-known luciferase. Well-known genetic engineering techniques can be used to introduce mutation into an amino acid sequence. In this case firstly, a mutation such as an addition, deletion, or substitution is introduced into a nucleotide sequence of a luciferase gene derived from the abovementioned luminescent organism or a well-known luciferase gene by genetic engineering techniques so as to generate a mutant luciferase gene. Subsequently, the mutant gene is incorporated into an appropriate host-vector system, thereby generating a recombinant microorganism. Then the recombinant microorganisms producing the luciferase of the present invention are selected by screening. The selected recombinant microorganisms are cultured in a medium. Finally the luciferase can be collected from the culture product.

Hereinafter surfactant-resistant luciferase obtained by introduction of a mutation into an amino acid sequence is referred to as "mutant luciferase."

The mutant luciferase is for example, luciferase wherein an amino acid corresponding to an amino acid at the 490-position of the GENJI firefly luciferase or the HEIKE firefly luciferase, is substituted by an amino acid other than glutamic acid in an amino acid sequence of a wild-type firefly luciferase. The amino acid other than glutamic acid is a basic amino acid. Specific examples include lysine, arginine, and histidine. The

term "an amino acid corresponding to the amino acid at the 490-position of the GENJI or the HEIKE firefly luciferase" means an amino acid corresponding to the amino acid at the 490-position of the GENJI or HEIKE firefly luciferase when the determined amino acid sequence of luciferase is compared to an amino acid sequence of the GENJI or HEIKE firefly luciferase.

Moreover, in the GENJI or HEIKE firefly luciferase, the amino acid at the 490-position is glutamic acid. Further, in North American firefly luciferase, "an amino acid corresponding to the amino acid at the 490-position of the GENJI or the HEIKE firefly luciferase" corresponds to the glutamic acid at the 487-position.

More specifically, the mutant luciferase is a polypeptide comprising an amino acid sequence shown in SEQ ID NO:1 or 2, or said amino acid sequence wherein one or more amino acids are added, deleted or substituted.

[Method for producing mutant luciferase by genetic engineering techniques]

A method for generating mutant luciferase by genetic engineering techniques will now be described as follows.

The mutant luciferase is produced by introducing mutation such as additions, deletions, and substitutions into a nucleotide sequence of known luciferase and allowing an appropriate vector-host system to express the gene.

The known luciferase genes includes, but are not limited to, a firefly luciferase gene, more specifically a wild-type HEIKE firefly luciferase gene (Japanese Patent Application Laid-Open No. 2-171189) and a thermostable HEIKE firefly luciferase gene (Japanese Patent Application Laid-Open No. 5-244942).

i) A method for introducing mutation into a luciferase gene is, for example a

method wherein the gene and a mutagen are allowed to contact with each other. Specific examples of the mutagen include hydroxylamine, nitrous acid, sulfurous acid, and 5-bromouracil. Further, ultra violet irradiation, cassette mutagenesis, and site-directed mutagenesis using PCR can also be used. Furthermore, a mutant lucefirase gene having a mutation at a desired position can be generated by annealing chemically synthesized DNA.

- ii) Next, the mutant luciferase gene is inserted into a vector DNA having such as a promoter sequence, a marker gene, and a replication origin, etc, thereby producing a recombinant plasmid. Any vector DNA can be used so far as it can be replicated in a host cell. Examples of the vector DNA include plasmid DNA and bacteriophage DNA. When the host cell is *Escherichia coli*, examples of the vector DNA include plasmid pUC119 (Takara Shuzo Co., Ltd.), pBluescript SK+(Stratagene), pMAL-C2 (NEW England Labs), pGEX-5X-1 (Pharmacia), pXa1 (Boehringer), and pMA56 (G.Ammerer, Meth. Enzymol., 101, 192, 1983).
- Subsequently, an appropriate host cell is transformed or transduced with the above recombinant plasmid, and screening is performed for recombinant microorganisms having the ability to produce the mutant luciferase.

Any host cells including eucaryotic and prokaryotic cells can be used. The eucaryotic cells include animal, plant, insect, yeast cells. The prokaryotic cells include *Escherichia coli*, *Bacillus subtilis*, and *Actinomyces*. The animal cells include CHO, COS, HeLa cells and cells of myeloma cell lines. The prokaryotic cells include microorganisms belong to the genus Escherichia, such as *Escherichia coli* JM101 (ATCC 33876), JM109 (produced by Takara Shuzo Co., Ltd.), XL1-Blue (produced by Stratagene), and HB101 (ATCC33694).

Transformation in the present invention can be performed by for example, D.M. Morrison's method (Meth. Enzymol., 68, 326-331, 1979); Transduction can be

conducted by for example, B.Hohn's method (Meth. Enzymol., 68, 299-309, 1979).

Methods for purification of recombinant DNA from recombinant microorganisms include P.Guerry's method (J.Bacteriology, 116, 1064-1066, 1973), and D.B.Clewell's method (J.Bacteriology, 110, 667-676, 1972).

The nucleotide sequence of a gene inserted into the recombinant DNA can be determined by, for example Maxam-Gilbert method (Proc. Natl. Acad. Sci. USA, 74, 560-564, 1977), and Dideoxy method (Proc. Natl. Acad. Sci. USA, 74, 5463-5467, 1977).

iv) The mutant luciferase of the present invention can be produced by culturing the recombinant microorganisms obtained in the manner described above in media.

When the host cell is *Escherichia coli*, recombinant E.coli may be cultured by solid culture methods, preferably liquid culture methods.

A culture medium of the present invention contains one or more nitrogen sources, such as yeast extract, tryptone, peptone, meat extract, corn steep liquor or exudate of soy bean or wheat bran, to which one or more of inorganic salts, such as sodium chloride, potassium phosphate, dipotassium phosphate, magnesium sulfate, magnesium chloride, ferric chloride, ferric sulfate or manganese sulfate are added. If necessary sugar and vitamins are added to this medium. Further the initial pH of the medium is preferably adjusted within pH 7 to 9. Moreover the culture is performed at a temperature within 30°C to 42°C, preferably at around 37°C for 3 to 24 hours, preferably for 5 to 8 hours. Preferable culture methods include aeration-agitation submerged culture, shaking culture, and static culture.

To recover mutant luciferase from the culture product after the completion of culturing recombinant E.coli, standard means for collecting enzymes can be employed. That is, the culture product is centrifuged to obtain cells. Then the cells are disrupted by treatment with lytic enzymes, such as lysozyme, ultrasonication, or milling. Fused

protein is discharged out of the cell. Subsequently insoluble substances are removed by filtration or centrifugation, so that a crude enzyme solution containing mutant luciferase can be obtained.

In the present invention the above crude enzyme solution can be used as authetic protein matter, or alternatively it can further be purified to higher purity by standard protein purification techniques. These techniques including sulfate salting out, organic solvent precipitation, ion exchange chromatography, hydrophobic chromatography, gel filtration chromatography, adsorption chromatography, affinity chromatography, and electrophoresis can be used solely or in combination.

The use of surfactant-resistant luciferase of the present invention allows the addition of a surfactant at a high concentration in the extraction process for intracellular ATP.

[Detection of intracellular ATP of the present invention]

Detection of intracellular ATP of the present invention will be described as follows.

i) First, ATP extraction reagent containing surfactant as an effective component is added to a sample containing cells so as to extract intracellular ATP out of the cells. The term "cells" refers to the cells derived from animal, plant, microorganism (e.g., yeasts, mold, fungi, bacteria, actinomyces, unicellular algae, viruses, and protozoa).

Any sample can be used so far as it contains the above cells. These samples include, but are not limited to, foods and drinks, pharmaceuticals, cosmetics, seawater, river water, industrial water, sewage, soil, urine, feces, blood, sputum, pus, and culture product of the above cells. A sample solution can also be prepared by suspending these samples in an appropriate solvent, such as distilled water, physiological saline, phosphoric acid buffer, Tris buffer, or sodium acetate

buffer. When a fluid specimen contains solids, the fluid specimen is suspended in an appropriate solvent or homogenized using a mixer so that it can be handled in the same manner as that in liquid form.

A sample of a filter membrane can also be prepared by filtering the above sample in liquid form through a hydrophilic or hydrophobic filter membrane. The hydrophilic or hydrophobic filter membrane by which cells are captured can be used as a sample. In such a case, a film- or sheet-type hydrophilic filter hydrophilic polytetrafluoroethylene, hydrophilic made of membrane acetylcellulose, and polyvinylidenefluoride, hydrophilic polyamide, nitrocellulose, etc., can be used. Hydrophobic filter membranes made of PVDF (polyvinylidenefluoride), PTFE (polytorafluoroethylene), and PE (polyethylene) etc., can be used.

Surfactants include anionic surfactants, cationic surfactants, ampholytic surfactants, and non-ionic surfactants.

Anionic sulfactants include sodium dodecyl sulfate (SDS), lauryl potassium sulfate, sodium monolauroyl phosphate, and sodium alkylbenzenesulfonic acid. Cationic surfactants include benzalkonium chloride (BAC), benzetonium chloride (BZC), cetylpyridinium chloride, cethyltrimethylammonium bromide, and myristyldimethylbenzylammonium chloride. ampholytic surfactants include Twittergent Detergent 3-08, 3-10, 3-12, 3-14, 3-16, and Tego. Finally non-ionic surfactants include Tween 20, 60, and 80, Span 60 and 80, Triton X-45 and x-100, polyoxyethylene ether, and polyoxyethylene lauryl ether.

Any concentration of a surfactant can be employed so far as it allows full expression of the ability to extract ATP. Preferable concentration of a surfactant is 0.05% or more of the mixture of a sample and ATP extraction reagent.

A sample and ATP extraction reagent are contacted with from each other at room temperature or with heating.

ii) After ATP extraction, bioluminescent reagent is added to the sample containing surfactant-resistant luciferase so as to cause emission. Then the light emission is measured.

When surfactant-resistant luciferase is derived from a firefly, the bioluminescent reagents are those containing e.g., the following components (a) to (c).

- (a) surfactant-resistant luciferase
- (b) luciferin
- (c) magnesium ions or other metal ions

Further in addition to the above components, substances involving pH preparation or improved shelf life may be added. Such substances include EDTA 2Na, dithiothreitol, ammonium sulfate, sucrose, 2-mercaptoethanol, HEPES, Tricine, and Tris.

iii) The amount of light emitted by the addition of a bioluminescent reagent can be measured by a luminometer such as a lumitester K-100 produced by Kikkoman Corporation, a luminescence reader BLR-201 produced by Aloka Co.,Ltd. (an improved type, or a Lumat LB9501 produced by Berthold. When a filter membrane by which cells are captured is used as a sample, the cells can be counted using a bioluminescent image analysis system device to photograph spots on the filter membrane. Such a device is ARGUS-50/CL (with taper fiber: produced by Hamamatsu Photonics K.K.).

The present invention will now be described in detail by the use of examples.

However the technical field of the present invention is not limited by these examples.

Example 1 Surfactant resistance of natural type luciferase derived from various

fire fly species.

(Method of preparing wild type luciferase derived from various firefly species)

Luciferase derived from GENJI and HEIKE fireflies was prepared according to the

1 mM ethylene diamine-4-acetate-2-sodium and 2mM following methods.

phenylmethylsulfonylfluoride were added to 25mM Tris (hydroxy) aminomethane-

hydrochloric acid buffer. Further ammonium sulfate was added to this solution so as

to achieve 10% saturation. Tail portions of the various firefly species were added to

this mixture at pH 7.8, and then disrupted using Hiskotoron (produced by

Nichionrikakikaiseisakusho). The resulting solution was centrifuged at 12,000 r.p.m.

for 20 minutes to obtain supernatants as starting materials for purification.

purification was conducted by the process comprising salting out of ammonium sulfate,

Ultrogel Ac A34 (produced by LKB) column, and hydroxy-apatite HPLC (produced

byTOSHOH, TSK gel HA-1000) column. Finally an electrophoretically homogenous

a commercial product (Sigma, L-9506).

(Method of determining luciferase activity)

A luciferase sample was properly diluted using enzyme-diluted solution (1mM EDTA,

1mM 2-mercaptoethanol, 1% BSA, 50mM HEPES, (pH7.5)). To $100 \,\mu$ l of this

solution, 100 µ1 of substrate solution (1.4mM luciferin, 40mM ATP, 300mM MgSO₄.

 $7H_2O$, 50mM HEPES, (pH 7.5)) was added.

The light emission was measured using BLE-201 Luminescence reader (produced by

Aloka Co., Ltd.) under the following conditions.

Measuring range: x100

Numerical value displayed: x1000

Measuring temperature: 30°C

Measuring time: 20 seconds

1MLU (mega light unit) /ml is a value for activity when the measured value under these conditions was 1 Kcount.

(Method of determining surfactant-resistance)

Enzyme samples were obtained by preparing luciferase samples derived from various firefly species using enzyme-diluted solution (1mM EDTA, 1mM 2-mercaptoethanol, 5% glycerol, 50mM HEPES, (pH7.5)) to achieve 0.5 MLU/ml concentration.

 $50\,\mu 1$ of 0.4% benzalkonium chloride (25mM Tricine at pH 7.75) and then $50\,\mu 1$ of the enzyme sample were added to 100 μ l of substrate solution (4mM ATP, 0.4mM luciferin, 10mM magnesium sulfate, 50mM HEPES (pH 7.5)). After the solution was stirred for 5 seconds, the light emission was measured every second using Berthold Lumat LB-9501 for 1 minute.

Figure 2 shows the results. Along the vertical axis in this figure, the relative ratio of the light emission was plotted with the initial amount of light emitted considered to be 100% upon use of 25mM Tricine (pH 7.75) instead of 0.4% benzalkonium chloride.

As shown in these results, North American firefly luciferase was low in the initial light emission and the light emission decayed rapidly. This was caused by the low surfactant-resistance of the North American firefly luciferase. This can lead to low sensitivity and accuracy in measuring such values. On the other hand, GENJI firefly luciferase showed an initial light emission higher than that of North American firefly luciferase. That is, GENJI firefly luciferase was shown to have a surfactant resistance superior to that of North American firefly luciferase. Furthermore, HEIKE firefly luciferase showed an initial light emission higher than that of GENJI firefly luciferase and the emission decayed slowly. Therefore, HEIKE firefly luciferase has good surfactant resistance, superior to that of GENJI firefly luciferase. These results suggest that the degree of surfactant resistance of luciferase varies according to the firefly species.

Example 2 Preparation of mutant luciferase HLK and HIK

Two types of mutant luciferase (named "HLK" and "HIK") were prepared according to the following methods.

(Production of a gene encoding mutant luciferase HLK)

A mutant luciferase gene was produced by site-directed mutagenesis using PCR. A plasmid pHLf7-217Leu described in Japanese Patent Application Laid-Open No. 5-244942 was used as a template for PCR reaction. The pHLf7-217Leu was a recombinant plasmid prepared by inserting a thermostable HEIKE firefly luciferase gene, in which an amino acid corresponding to Ala at the 217-position was substituted for a Leu-encoding gene, into a plasmid pUC119. In addition, E. coli JM101, to which the recombinant plasmid pHLf7-217Leu had been introduced, has been named E.coli JM101 (pHLf7-217Leu) and was deposited on April 22, 1992 as FERM BP-3840 with the National Institute of Bioscience and Human-Technology, Agency of Industrial Science and Technology (1-3, Higashi 1-chome, Tsukuba-shi, Ibaraki-ken, Japan).

The primer for PCR reaction was an oligonuceltide having a nucleotide sequence shown in SEQ ID No: 1 or 2. The DNA polymerase was a KOD dash polymerase (produced by TOYOBO). A PCR reaction cycle (94°C for 30 seconds, 50°C for 2 seconds, and 74°C for 3 minutes) was repeated for 30 times according to the examples attached to KOD dash polymerase. The PCR product was ligated into a circular recombinant plasmid pHLfLK using standard techniques.

Sequencing of a mutant luciferase gene contained in the pHLfLK was performed.

Reaction was conducted using a Diprimer Taq Sequencing Kit (produced by Applied Biosystems). Then the eletrophoretic analysis was performed using ABI 373A DNA sequencer (produced by Applied Biosystems). The entire nucleotide sequence of the obtained mutant luciferase gene is shown in SEQ ID NO: 3, and the amino acid sequence of a polypeptide encoded by this gene is shown in SEQ ID NO: 4. In the mutant luciferase gene, the genetic portion corresponding to alanine at the 217-position of wild-type HEIKE firefly luciferase was substituted by a gene encoding leucine, the genetic portion corresponding to glutamic acid at the 490-position of the same was substituted by a gene encoding lysine. The pHLfLK-introduced E.coli JM109 strain was named E.coli JM109 (pHLfLK) (see Figure 1). E. coli JM109 (pHLfLK) was deposited as FERM BP-6147 on October 16, 1997) with the National Institute of Bioscience and Human-Technology, Agency of Industrial Science and Technology, Japan.

The polypeptide shown in SEQ ID NO:4 was named the mutant luciferase HLK.

(Preparation of gene encoding mutant luciferase HIK)

A mutant luciferase gene was prepared using the plasmid pHLf7-217Ile described in Japanese Patent Application Laid-Open No. 5-244942. The plasmid pHLf7-217Ile was a recombinant plasmid prepared by inserting a thermostable HEIKE firefly luciferase gene, in which an amino acid corresponding to Ala at the 217-position was substituted for a Ile-encoding gene, into a plasmid pUC119. The transformant strain obtained using this plasmid was deposited on April 22, 1992 as FERM BP-3841 with the National Institute of Bioscience and Human-Technology, Agency of Industrial Science and Technology, Japan.

About a 560bp fragment obtained by cutting the pHLfLK with EcoRV and NarI was obtained by agarose gel electrophoresis. Then the fragment was inserted into the pHLf7-217Ile treated with the same restriction enzymes.

The resulting recombinant plasmid has been named pHLfIK and the plasmid-introduced E.coli JM109 strain has been named E.coli JM109 (pHLfIK).

E.coli JM109 (pHLfIK) was deposited on October 16, 1997 as FERM BP-6146 with the National Institute of Bioscience and Human-Technology, Agency of Industrial Science and Technology, Japan.

The entire nucleotide sequence of the mutant luciferase gene contained in the pHLfIK is shown in SEQ ID NO: 5, and the amino acid sequence of a polypeptide encoded by this gene is shown in SEQ ID NO: 6. In the mutant luciferase gene, the genetic portion corresponding to alanine at the 217-position of wild-type HEIKE firefly luciferase was substituted by a gene encoding isoleucine, the genetic portion corresponding to glutamic acid at the 490-position of the same was substituted by a gene encoding lysine (see Fig. 1).

A polypeptide shown in SEQ ID NO:6 was named the mutant HIK firefly.

Example 3 Preparation of mutant luciferase HLK and HIK

E.coli JM109 (pHLfLK) and E.coli JM109 (pHLfIK) were inoculated on LB media (1% Bacto-trypton (W/V), 0.5% yeast extract (W/V), 0.5% NaCl (W/V), ampicillin (50 μ g/ml), 1.4% agar (W/V)), each containing ampicillin, and cultured at 37°C for 18 hours. The resulting culture fluid was centrifuged at 8000 r.p.m. for 10 minutes. The precipitated cells were suspended in 0.1M potassium phosphate buffer at pH 7.8 (0.1M ammonium sulfate, 1mM EDTA) were disrupted by ultrasonication.

Next, crude enzyme solution was obtained by centrifugation at 12000 r.p.m. for 10 minutes. The obtained enzyme solution was purified using the above purification techniques such that it becomes an electrophoretically homogenous sample.

Example 4 Surfactant resistance of mutant luciferase HLK and HIK

(Changes in emission with time)

To compare surfactant resistance of mutant luciferase with that of known luciferase, changes in emission with time were measured according to the aforementioned methods of measuring surfactant resistance. Figure 3 shows the results obtained by the use of 0.4% benzalkonium chloride (25mM Tricine (pH 7.75)). Figure 4 shows the results obtained by the use of 0.8% benzethonium chloride (25mM Tricine (pH 7.75)).

"HEIKE I mutant" in this figure is thermostable HEIKE firefly luciferase (described in Japanese Patent Application Laid-Open No. 5-244942) wherein Ala at the 217-position of wild-type HEIKE firefly luciferase is substituted for Ile. "HEIKE L mutant" is thermostable HEIKE firefly luciferase (Japanese Patent Application Laid-Open No. 5-244942) wherein Ala at the 217 position of wild-type HEIKE luciferase is substituted by Leu. "HIK" is a mutant wherein Glu at the 490-position of HEIKE I mutant is substituted by Lys, that is, the mutant luciferase HIK prepared in Example 3. "HLK" is a mutant wherein Glu at the 490-position of HEIKE L mutant is substituted by Lys, that is, the mutant luciferase HLK prepared in Example 3.

As can be seen in Fig. 3 which shows the results for benzalkonium chloride, the emission of HIK decayed more slowly than that of the HEIKE I mutant. Comparison of HLK and HEIKE L mutant reveals that HLK had initial light emission improved by about 20%, and slower decay in the light emission.

Therefore, the substitution of an amino acid at the 490-position resulted in improved surfactant-resistance of a luciferase.

As shown in Fig. 4 which shows the results obtained by the use of benzethonium chloride, HIK showed decay in emission more slowly than that of HEIKE I mutant. Further HLK showed slower decay in light emission than that of HEIKE L mutant. Therefore, the substitution of an amino acid at the 490-position resulted in improved

surfactant resistance.

(Comparison of emission rate)

The influence of the enzyme solution, substrate solution and benzalkonium chloride used when measuring change with time, on the measurement values taken under actual emission measurement conditions, was examined. Table 1 shows the light emission measured using Berthold Lumat LB-9501 under measuring conditions (5 seconds of waiting time, 3 seconds of measuring time).

In addition, the emission rate (remaining activity) was calculated by dividing the light emission measured in the presence of 0.4% benzalkonium chloride by a control value. Here the control value was the light emission upon use of 25mM Tricine at pH 7.75 instead of 0.4% benzalkonium chloride.

Table 1

Luciferase type	Light emissi	Emission rate (%)	
	Without extraction	With extraction	
	reagent	reagent	
North American firefly	452563	97790	21.6
GENJI firefly	409406	167805	41.0
HEIKE firefly	425792	324724	76.3
HEIKE I mutant	422269	341039	80.8
HEIKE L mutant	423728	343634	81.1
HIK	386429	345159	89.3
HLK	390289	396764	101.7

North American firefly luciferase showed an emission rate as low as 21.6%, suggesting a large decrease in sensitivity. On the other hand, the emission rates for GENJI and HEIKE firefly luciferase were 41.0% and 76.3%, respectively, suggesting that the sensitivity of these firefly luciferases were less affected than that of North American firefly luciferase.

The emission rate for mutant luciferase HIK and HLK were 89.3% and 101.7%,

respectively. These rates were far greater than those of wild-type HEIKE firefly luciferase and thermostable HEIKE firefly luciferase. Particularly the emission rate of HLK was almost 100%. That is, HLK can yield the same light emission regardless of the presence or absence of a surfactant. Therefore, the sensitivity of HLK is totally unaffected by the use of a surfactant, allowing measurement with high accuracy.

(Comparison of IC50)

Benzalkonium chloride and various luciferases were contacted with each other for 10 minutes. Then the benzalkonium chloride concentration (IC50), at which activity is inactivated by 50% was determined. Equal amounts of luciferase solution prepared at this concentration and 0.01 to 0.1% benzalkonium chloride were mixed, and then allowed to stand for 10 minutes at room temperature. Subsequently, $100 \,\mu$ 1 of substrate solution was added to the mixture. Immediately after addition, the light emission was measured using Berthold Lumat LB-9501. IC50s obtained were as shown in Table 2.

Table 2 IC_{50} for various luciferase

• •	
Luciferase type	IC ₅₀ (%)
North American firefly	0.014
GENJI firefly luciferase	0.016
HEIKE firefly luciferase	0.026
HEIKE I mutant	0.028
HEIKE L mutant	0.028
HIK	0.032
HLK	0.035

North American firefly luciferase showed the lowest IC_{50} among the three types of wild-type luciferase. That is, North American firefly luciferase was shown to have the lowest resistance to a surfactant. HEIKE firefly luciferase showed the highest IC_{50} among the wild-type luciferase. HLK and HIK showed IC_{50} higher than those of wild-type HEIKE firefly luciferase and thermostable HEIKE firefly luciferase, suggesting that the resistance was improved by the substitution of an amino acid at the 490-position.

Especially HLK showed IC₅₀ higher than that of HIK, indicating that HLK possesses the best surfactant-resistance.

Example 5 Method for measuring intracellular ATP

Next, a method for measuring intracellular ATP using the surfactant-resistant luciferase of the present invention will be described.

A standard technique used herein was TCA extraction method wherein intracellular ATP is extracted using trichloroacetic acid (TCA) and the amount of ATP extracted is measured using luciferin-luciferase luminescence reaction. TCA extraction method is excellent in extraction efficiency. Further in TCA extraction method no inhibition of luminescence reaction is caused by TCA because emission is measured after the sample containing TCA is diluted 1:100. Because of this dilution, however, TCA extraction method is complicated and can cause a decrease in the measuring sensitivity.

1. Materials

(1) Surfactant

Benzalkonium chloride (BAC, Japanese Pharmacopoeia) was used. ATP extraction reagent was prepared by dissolving this surfactant at 0.25% concentration into 25mM Tricine (pH 7.75).

(2) Microorganisms

Four strains, Escherichia coli (ATCC 25922), Staphylococcus aureus (ATCC 25923), Pseudomonas aeruginosa (ATCC 27853) and Enterococcus faecalis (ATCC 29212) were used.

(3) Preparation of samples

In standard techniques, a sample, undiluted solution, was prepared by culturing the prescribed microorganisms on a normal broth medium (produced by Eiken chemical Co., Ltd.) at 35°C overnight. In the method of the present invention, a sample diluted solution was prepared by diluting an undiluted solution of the culture fluid to 1:100 with

sterile water.

(4) Luciferase

Surfactant-resistant luciferase of the present invention were HIK and HLK. Control surfactant-resistant luciferase types were known luciferase (North American firefly luciferase, GENJI firefly luciferase, HEIKE firefly luciferase, HEIKE I mutant, and HEIKE L mutant).

(5) Luminescence reagent

Luminescence reagent was prepared by adding various luciferase to solution containing 0.15mM luciferin, 6mM EDTA, 15mM magnesium acetate, 0.2mM dithiothreitol, 0.5% BSA and 25mM HEPES (pH 7.75).

The amount of luciferase to be added was prepared such that the light emission produced when $100\,\mu\,l$ of $2x10^{-8}$ M ATP standard solution was added to $100\,\mu\,l$ of the luminescence reagent would be the same amount of the light emission produced when a luminescence reagent attached to Luciferase LU (Kikkoman Corporation) was used.

2. Method for measuring intracellular ATP

(1) Method of the present invention

ATP extraction reagent $100\,\mu$ l was added to $100\,\mu$ l of a sample. The solution was allowed to stand for 20 seconds at room temperature. Then $100\,\mu$ l of the luminescence reagent was added to this solution. Immediately after addition, the light emission was measured using Lumat LB-9501 produced by Berthold.

(2) Standard technique

10 % trichloro acetate solution $100\,\mu\,\mathrm{l}$ was added to $100\,\mu\,\mathrm{l}$ of a sample and the solution was allowed to stand for 1 minute. 25 mM Tricine (pH 7.75) 9.8ml was added to this extract, and then the extract was well stirred. 25 mM Tricine (pH 7.75) and $100\,\mu\,\mathrm{l}$ of a luminescence reagent attached to CheckLite LU (produced by Kikkoman Corporation) were added to $100\,\mathrm{l}$ of the sample. Immediately after addition, the light emission was measured using Lumat LB-9501 produced by Berthold.

3. Results

Tables 3 and 4 show the results. The relative ratio of the light emissions obtained by the use of the luminescence reagents using various luciferase types is also shown in these tables. Here the light emission obtained by the standard technique (TCA extraction method) was defined as 100%.

Table 3 Detection of intracellular ATP

Measuring method	E.coli ATCC25922		S.aureus ATCC 25923	
	Measured value (RLU)	Relative ratio (%)	Measured value(RLU)	Relative ratio (%)
Standard technique (TCA extraction method)	132794	(100.0)	130220	(100.0)
North American firefly	153	(0.1)	163	(0.1)
GENJI firefly	463	(0.3)	659	(0.5)
luciferase				
HEIKE firefly	76082	(57.3)	74019-	(56.8)
luciferase				
HEIKE I mutant	47655	(35.9)	50031	(38.4)
HEIKE L mutant	46217	(34.8)	51243	(39.4)
HIK	97073	(73.1)	76533	(58.8)
HLK	87981	(66.3)	72182	(55.4)

Table 4 Detection of intracellular ATP

Measuring method	P.aeruginosa ATCC 27853		E.faecalis ATCC 29212	
	Measured value (RLU)	Relative ratio (%)	Measured value(RLU)	Relative ratio (%)
Standard technique (TCA extraction method)	168141	(100.0)	12427	(100.0)
North American firefly	553	(0.3)	113	(0.1)
GENJI firefly luciferase	1503	(0.9)	163	(1.3)
HEIKE firefly luciferase	117096	(69.6)	8132	(65.4)
HEIKE I mutant	80455	(47.8)	4586	(36.9)
HEIKE L mutant	81069	(48.2)	4762	(38.3)
HIK	131134	(78.0)	7914	(63.7)
HLK	131815	(78.4)	7998	(64.4)

No emission was observed for the luminescence reagent containing North American firefly luciferase. GENJI firefly luciferase showed weak emission. This is because the luciferase itself was devitalized by the surfactant. Therefore, it was shown that the surfactant at high concentration such as was used in this examination cannot be used as an ATP extraction reagent for the luciferase.

Unlike North American firefly luciferase and GENJI firefly luciferase, HEIKE firefly luciferase showed emission 60 to 70% of that in TCA extraction method. HEIKE firefly luciferase was shown to possess surfactant-resistance higher than those of North American firefly luciferase and GENJI firefly luciferase.

Light emissions from HEIKE L mutant, and HEIKE I mutant which is thermostable HEIKE firefly luciferase were each equivalent to around 40% of that in TCA extraction method, and largely lower than that of wild-type HEIKE firefly luciferase.

Each of the light emission from HIK and HLK, which is surfactant-resistant luciferase

of the present invention, respectively was more intense than that from wild-type HEIKE

luciferase and thermostable luciferase. Further the light emission in this case was

equivalent to 60 to 80% of that in TCA extraction method.

HIK and HLK are mutants wherein Glu at the 490-position of HEIKE I and HEIKE L

mutants are substituted for Lys, respectively. That is, the introduction of said mutation

into the amino acid at the 490-position improved resistance to a surfactant. The

sensitivity of HIK and HLK is less affected by ATP extraction reagent even at such a

high concentration employed in this examination, suggesting the use of HIK and HLK

enable highly accurate measurement.

Industrial Applicability

The use of a novel surfactant-resistant luciferase according to the present invention for

measuring intracellular ATP allows the detection without a decrease in luciferase

activity even in the presence of a surfactant at a high concentration.

All publications, patents and patent applications cited herein are incorporated herein by

reference in their entirety.

Sequence Listing Free Text

SEQ ID NO:1: A synthetic DNA

SEQ ID NO:2: A synthetic DNA

25

CLAIMS

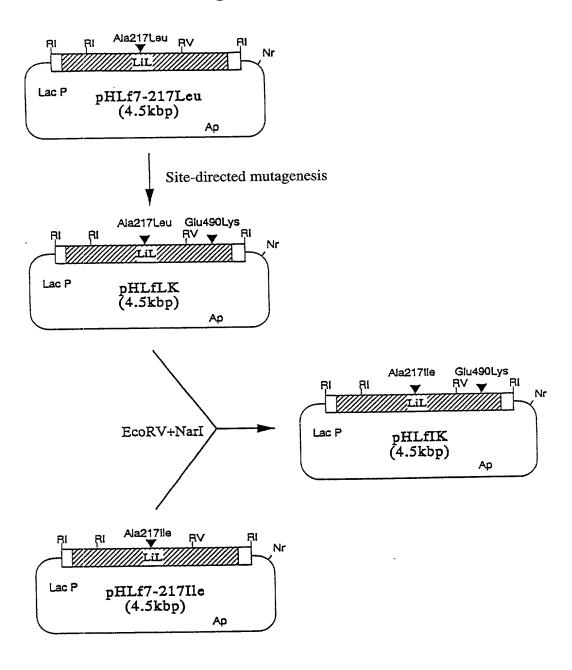
- 1. A luciferase having resistance to a surfactant.
- 2. The luciferase of claim 1 wherein an amino acid corresponding to that at the 490-position of luciferase from Genji or Heike firefly is substituted by an amino acid other than glutamic acid in the amino acid sequence of firefly luciferase.
- 3. The luciferase of claim 2 wherein the amino acid other than glutamic acid is lysine.
- 4. The luciferase of claim 1 wherein it is:
 - (a) a polypeptide consisting of the amino acid sequence shown in SEQ ID NO:4; or
 - (b) a polypeptide comprising additions, deletions or substitutions of one or more amino acids in the amino acid sequence of the polypeptide defined in (a) and having luciferase activity resistant to a surfactant.
- 5. The luciferase of claim 1 wherein it is:
 - (a) a polypeptide consisting of the amino acid sequence shown in SEQ ID NO:6; or
 - (b) a polypeptide comprising additions, deletions, or substitutions of one or more amino acids in the polypeptide defind in (a) and having luciferase activity resistant to a surfactant.
- 6. A luciferase gene encoding the luciferase of any one of claim 1 to 5.
- 7. A recombinant vector comprising the luciferase gene of claim 6.
- 8. A transformant comprising the recombinant vector of claim 7.
- 9. A method for producing a luciferase wherein the method comprising culturing the transformant of claim 8 in a medium and recovering the luciferase from the resulting culture.
- 10. A method for measuring intracellular ATP characterized in that a luciferase having resistance to a surfactant is used as a luciferase for use in the method comprising a first step wherein ATP is extracted in the presence of the surfactant from cells in a sample, a second step wherein a luminescence reagent containing luciferase is added to the extracted ATP solution to cause emission of light, and a third step wherein the amount of light emission is measured.

- 11. The method for measuring intracellular ATP of claim 10 wherein the luciferase having resistance to a surfactant is a luciferase of any one of claim 1 to 5.
- 12. The method for measuring intracellular ATP of claim 10 or 11 wherein the light emission is caused by addition of a luminescence reagent in the presence of a surfactant of 0.01% or more.
- 13. The method for measuring intracellular ATP of claim 10, 11 or 12 wherein the surfactant is any of a cationic surfactant, an anionic surfactant, a nonionic surfactant, and a ampholytic surfactant.

Abstract

The present invention relates to luciferase having resistance to a surfactant and a method for measuring intracellular ATP which is characterized in that the luciferase having resistance to a surfactant is used in this method comprising the steps of: a first step wherein ATP is extracted from cells in a sample; a second step wherein light emission is produced by adding a luminescence reagent containing luciferase to the extracted ATP solution; and a third step wherein the light emission is measured.

Figure 1



LIL; Luciola lateralis luciferase cDNA, Ap; β -lactamase gene, LacP; β -galactosidase promoter, RI; EcoRI, RV;EcoRV, Nr;Narl

Figure 2

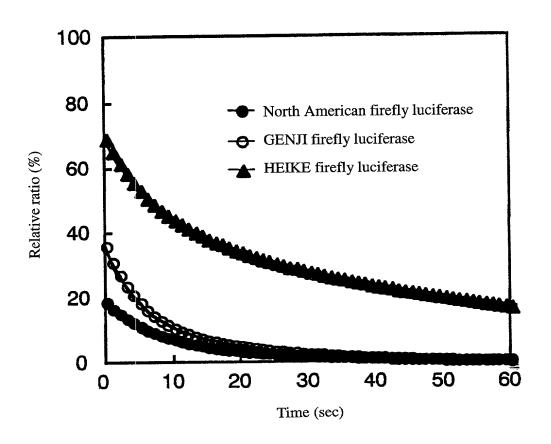
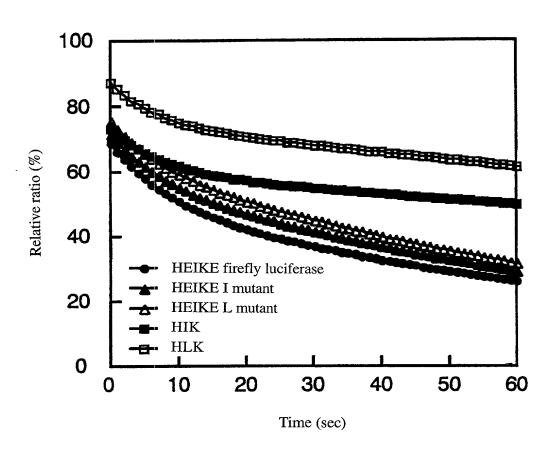


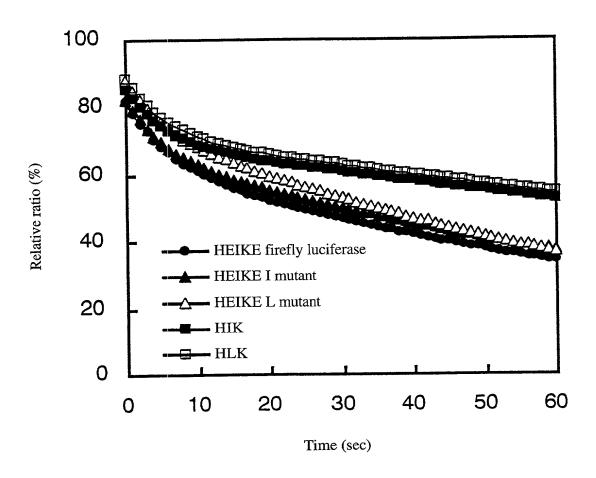
Figure 3



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Figure 4



Attorney's Docket	No.:

DECLARATION, POWER OF ATTORNEY AND PETITION

I (We), the undersigned inventor(s), hereby declare that: My residence, post office address and citizenship are as stated below next to my name, I (We) believe that I am (we are) the original, first, and joint (sole) inventor(s) of the subject matter which is claimed and for which a patent is sought on the invention entitled LUCIFERASE AND METHODS FOR MEASURING INTRACELLULAR ATP USING THE SAME the specification of which is attached hereto. was filed on _____as Application Serial No. and amended on _____ was filed as PCT international application Number PCT/JP98/05864 on December 24, 1998 and was amended under PCT Article 19

I (We) hereby state that I (We) have reviewed and understand the contents of the above-identified specification, including the claims, as amended by any amendment referred to above; that I (We) do not know and do not believe that this invention was ever known or used before my invention or discovery thereof, or patented or described in any printed publication in any country before my invention or discovery thereof, or more than one year prior to this application, or in public use or on sale in the United States for more than one year prior to this application; that this invention or discovery has not been patented or made the subject of an inventor's certificate in any country foreign to the United States on an application filed by me or my legal representatives or assigns more than twelve months before this application.

on_____(if applicable).

I (We) acknowledge the duty to disclose information known to be material to the patentability of this application as defined in Section 1.56 of Title 37 Code of Federal Regulations.

I (We) hereby claim foreign priority benefits under Section 119(a)-(d) of Title 35 United States Code, of any foreign application(s) for patent or inventor's certificate listed below and have also identified below any foreign application for patent or inventor's certificate having a filing date before that of the application on which priority is claimed:

Priority

			11011	* 3	
Application No.	Country	Filing date	claime	ed	
361022/1997	Japan	December 26, 1997	Yes	\square No	
			☐ Yes	\square No	
			☐ Yes	□ No	
			☐ Yes	□ No	
of any United S	tates application(s)	listed below.			
(Application Nu	umber)	(Filing Date)			
(Application N	umber)	(Filing Date)			

I (We) hereby claim the benefit under Section 120 of Title 35 United States Code, of any United States application(s) listed below and, insofar as the subject matter of each of the claims of this application is not disclosed in the prior United States application in the manner provided by the first paragraph of Section 112 of Title 35 United States Code, I (We) acknowledge the duty to disclose material information as defined in Section 1.56(a) of Title 37 Code of Federal Regulations, which occurred between the filing date of the prior application and national or PCT international filing date of this application:

Application	Serial No.	Filing Date	patented, a b an d o n e d)

Status (pending,

And I (We) hereby appoint: Norman F. Oblon, Registration No. 24,618; Marvin J. Spivak, Registration No. 24,913; C. Irvin McClelland, Registration No. 21,124; Gregory J. Maier, Registration No. 25,599; Arthur I. Neustadt, Registration No. 24,854; Richard D. Kelly, Registration No. 27,757; James D. Hamilton, Registration No. 28,421; Eckhard H. Kuesters, Registration No. 28,870; Robert T. Pous, Registration No. 29,099; Charles L. Gholz, Registration No. 26,395; Vincent J. Sunderdick, Registration No. 29,004; William E. Beaumont, Registration No. 30,996; Robert F. Gnuse, Registration No. 27,295; Jean-Paul Lavalleye, Registration No. 31,451; Stephen G. Baxter, Registration No. 32,884; Martin M. Zoltick, Registration No. 35, 745; Robert W. Hahl, Registration No. 33,893; Richard L. Treanor, Registration No. 36, 379; Steven P. Weihrouch, Registration No. 32, 829; John T. Goolkasian, Registration No. 26, 142; Richard L. Chinn, Registration No. 34, 305; Steven E. Lipman, Registration No. 30, 011; Carl E. Schlier, Registration No. 34, 426; James J. Kulbaski, Registration No. 34, 648; Richard A Neifeld, Registration No. 35, 299; J. Derek Mason, Registration No. 35, 270; Surinder Sachar 34, 423; Christina M. Gadiano, Registration No. 37, 628; Jeffrey B. McIntyre, Registration No. 36, 867; and Paul E. Rauch, Registration No. 38, 591; our (my) attorneys, with full powers of substitution and revocation, to prosecute this application and to transact all business in the Patent Office connected therewith; and we (I) hereby request that all correspondence regarding this application be sent to the firm of OBLON, SPIVAK, McCLELLAND, MAIER & NEUSTADT, P.C. whose Post office address is: Fourth Floor, 1755 Jefferson Davis Highway, Arlington, Virginia 22202 U.S.A. I (We) declare further that all statements made herein of my (our) knowledge are true and that all statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

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09/581241 533 Rec'd PCT/PTO 26 JUN 2000

SEQUENCE LISTING

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0.00	an t	000	0++	+ o +	~~~	000	000	a + +	t 0.0	0.00	~~~	0.00	an t	0 + +	t t o	790
	gat															720
	Asp	riu	116	1 ) 1		ASII	GIII	Val	Sei		GIY	1111	НІА	116		
225					230					235					240	
ant	a to	ata	cen	t + 0	na t	co t	aa+	+++	nn+	atr	+++	2 C f	20+	tta	aac	769
acı	gta	gıa	cca	ııc	ual	cal	ggı	iii	ggı	aig	ııı	acı	ati	ııa	RRC	768

tat cta act tgt ggt ttt cgt att gtc atg tta acg aaa tt Tyr Leu Thr Cys Gly Phe Arg Ile Val Met Leu Thr Lys Ph	255 t gac gaa 816
	t gac gaa 816
	i gat gaa oro
Tyr Leu Thr Cys Gly Phe Arg Ile Val Met Leu Thr Lys Ph	_
	e Asp Glu
260 265 27	0
gag act ttt tta aaa aca ctg caa gat tac aaa tgt tca ag	c gtt att 864
Glu Thr Phe Leu Lys Thr Leu Gln Asp Tyr Lys Cys Ser Se	r Val Ile
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ctt gta ccg act ttg ttt gca att ctt aat aga agt gaa tt	a ctc gat 912
Leu Val Pro Thr Leu Phe Ala Ile Leu Asn Arg Ser Glu Le	u Leu Asp
290 295 300	
aaa tat gat tta tca aat tta gtt gaa att gca tct ggc gg	a gca cct 960
Lys Tyr Asp Leu Ser Asn Leu Val Glu Ile Ala Ser Gly Gl	y Ala Pro
305 310 315	320
tta tct aaa gaa att ggt gaa gct gtt gct aga cgt ttt aa	t tta ccg 1008
Leu Ser Lys Glu Ile Gly Glu Ala Val Ala Arg Arg Phe As	n Leu Pro
325 330	335
ggt gtt cgt caa ggc tat ggt tta aca gaa aca acc tct gc	a att att 1056
Gly Val Arg Gln Gly Tyr Gly Leu Thr Glu Thr Thr Ser Al	a Ile Ile
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340   345   35	
340 345 35	

lle Thr Pro Glu Gly Asp Asp Lys Pro Gly Ala Ser Gly Lys Val Val

355 360 365

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Pro	Leu	Phe	Lys	Ala	Lys	Val	Ile	Asp	Leu	Asp	Thr	Lys	Lys	Thr	Leu	
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ggc	ccg	aac	aga	cgt	gga	gaa	gtt	tgt	gta	aag	ggt	cct	atg	ctt	atg	1200
Gly	Pro	Asn	Arg	Arg	Gly	Glu	Val	Cys	Val	Lys	Gly	Pro	Met	Leu	Met	
385					390					395					400	
aaa	ggt	tat	gta	gat	aat	cca	gaa	gca	aca	aga	gaa	atc	ata	gat	gaa	1248
Lys	Gly	Tyr	Val	Asp	Asn	Pro	Glu	Ala	Thr	Arg	Glu	Ile	He	Asp	Glu	
				405					410					415		
	ggt															1296
Glu	Gly	Trp		His	Thr	Gly	Asp		Gly	Tyr	Tyr	Asp	Glu	Glu	Lys	
			420					425					430			
	ttc															1344
His	Phe		He	Vai	Asp	Arg		Lys	Ser	Leu	He		Tyr	Lys	Gly	
		435					440					445				
4-4		~+~		4			44.		1.1	-: <b>1 1</b>	- 4 4					1000
	caa															1392
ГУГ	Gln	vai	Pro	Pro	Ala		Leu	6111	ser	vai		Leu	GIN	HIS	Pro	
	450					455					460					
00+	0++	+++	go t	~~~	~~~	~ + +	a a t	~~~	~ + +	220	an t	a a t	0 + 0	~ o t	~~+	1 4 4 0
	att															1440
	lle	rne	дзр	ніа		Val	ніа	GIY	Val		АЅР	110	116	ына	-	
465					470					475					480	

1488 gag ctt ccg gga gct gtt gtt gta ctt aag aaa gga aaa tct atg act Glu Leu Pro Gly Ala Val Val Leu Lys Lys Gly Lys Ser Met Thr 490 485 495 gaa aaa gaa gta atg gat tac gtt gct agt caa gtt tca aat gca aaa 1536 Glu Lys Glu Val Met Asp Tyr Val Ala Ser Gln Val Ser Asn Ala Lys 500 505 510 cgt ttg cgt ggt ggt gtc cgt ttt gtg gac gaa gta cct aaa ggt ctc 1584 Arg Leu Arg Gly Gly Val Arg Phe Val Asp Glu Val Pro Lys Gly Leu 520 525 515 act ggt aaa att gac ggt aaa gca att aga gaa ata ctg aag aaa cca 1632 Thr Gly Lys Ile Asp Gly Lys Ala Ile Arg Glu Ile Leu Lys Lys Pro 530 535 540 gtt gct aag atg 1644 Val Ala Lys Met 545

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Met Asp Arg Tyr Ala Lys Leu Gly Ala Ile Ala Phe Thr Asn Ala Leu
35 40 45

Thr Gly Val Asp Tyr Thr Tyr Ala Glu Tyr Leu Glu Lys Ser Cys Cys
50 55 60

Leu Gly Glu Ala Leu Lys Asn Tyr Gly Leu Val Val Asp Gly Arg Ile
65 70 75 80

Ala Leu Cys Ser Giu Asn Cys Giu Giu Phe Phe Ile Pro Val Leu Ala 85 90 95

Gly Leu Phe Ile Gly Val Gly Val Ala Pro Thr Asn Glu Ile Tyr Thr
100 105 110

Leu Arg Glu Leu Val His Ser Leu Gly Ile Ser Lys Pro Thr Ile Val 115 120 125

Phe Ser Ser Lys Lys Gly Leu Asp Lys Val Ile Thr Val Gln Lys Thr
130 135 140

Val Thr Ala Ile Lys Thr Ile Val Ile Leu Asp Ser Lys Val Asp Tyr 145 150 155 160

Arg Gly Tyr Gln Ser Met Asp Asn Phe IIe Lys Lys Asn Thr Pro Gln

Gly Phe Lys Gly Ser Ser Phe Lys Thr Val Glu Val Asn Arg Lys Glu
180 185 190

Gln Val Ala Leu Ile Met Asn Ser Ser Gly Ser Thr Gly Leu Pro Lys 195 200 205

Gly Val Gln Leu Thr His Glu Asn Leu Val Thr Arg Phe Ser His Ala 210 215 220

Arg Asp Pro Ile Tyr Gly Asn Gln Val Ser Pro Gly Thr Ala Ile Leu 225 230 235 240

Thr Val Val Pro Phe His His Gly Phe Gly Met Phe Thr Thr Leu Gly
245 250 255

Tyr Leu Thr Cys Gly Phe Arg IIe Val Met Leu Thr Lys Phe Asp Glu 260 265 270

Glu Thr Phe Leu Lys Thr Leu Gln Asp Tyr Lys Cys Ser Ser Val Ile 275 280 285

Leu Val Pro Thr Leu Phe Ala Ile Leu Asn Arg Ser Glu Leu Leu Asp 290 295 300

Lys Tyr Asp Leu Ser Asn Leu Val Glu Ile Ala Ser Gly Gly Ala Pro 305 310 315 320 Leu Ser Lys Glu Ile Gly Glu Ala Val Ala Arg Arg Phe Asn Leu Pro 325 330 335

Gly Val Arg Gln Gly Tyr Gly Leu Thr Glu Thr Thr Ser Ala Ile Ile 340 345 350

Ile Thr Pro Glu Gly Asp Asp Lys Pro Gly Ala Ser Gly Lys Val Val
355 360 365

Pro Leu Phe Lys Ala Lys Val IIe Asp Leu Asp Thr Lys Lys Thr Leu 370 375 380

Gly Pro Asn Arg Arg Gly Glu Val Cys Val Lys Gly Pro Met Leu Met 385 390 395 400

Lys Gly Tyr Val Asp Asn Pro Glu Ala Thr Arg Glu IIe IIe Asp Glu
405 410 415

Glu Gly Trp Leu His Thr Gly Asp Ile Gly Tyr Tyr Asp Glu Glu Lys
420 425 430

His Phe Phe Ile Val Asp Arg Leu Lys Ser Leu Ile Lys Tyr Lys Gly
435
440
445

Tyr Gln Val Pro Pro Ala Glu Leu Glu Ser Val Leu Leu Gln His Pro
450 455 460

Asn Ile Phe Asp Ala Gly Val Ala Gly Val Pro Asp Pro Ile Ala Gly
465 470 475 480

Glu Leu Pro Gly Ala Val Val Leu Lys Lys Gly Lys Ser Met Thr 485 490 495

Glu Lys Glu Val Met Asp Tyr Val Ala Ser Gln Val Ser Asn Ala Lys
500 505 510

Arg Leu Arg Gly Gly Val Arg Phe Val Asp Glu Val Pro Lys Gly Leu 515 520 525

Thr Gly Lys Ile Asp Gly Lys Ala Ile Arg Glu Ile Leu Lys Lys Pro
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Val Ala Lys Met

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<220>

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	Tyr	Lys	Arg	Leu	Gln	Ala	Gly	Ala	Ser	Gly	Glu	Glu	Ile	Pro	Tyr	Phe
			30					25					20			
144	ctt	gca	aac	act	ttt	gct	att	gca	gga	ctt	aaa	gca	tat	cga	gat	atg
	Leu	Ala	Asn	Thr	Phe	Ala	Ile	Ala	Gly	Leu	Lys	Ala	Tyr	Arg	Asp	Met
				45					40					35		
192	tgt	tgc	tca	aaa	gaa	tta	tac	gaa	gcc	tac	acg	tat	gat	gtc	ggt	acc
	Cys	Cys	Ser	Lys	Glu	Leu	Tyr	Glu	Ala	Tyr	Thr	Tyr	Asp	Val	Gly	Thr
					60					55					50	
240	att	aga	gga	gat	gtt	gtt	ttg	ggt	tat	aat	aag	tta	gct	gag	gga	cta
	He	Arg	Gly	Asp	Val	Val	Leu	Gly	Tyr	Asn	Lys	Leu	Ala	Glu	Gly	Leu
	80					75					70					65
288	gcc	tta	gta	cct	att	ttt	ttc	gaa	gaa	tgt	aac	gaa	agt	tgc	tta	gcg
	Ala	Leu	Val	Pro	Пе	Phe	Phe	Glu	Glu	Cys	Asn	Glu	Ser	Cys	Leu	Ala
		95					90					85				
336	act	tac	att	gag	aat	act	cca	gct	gtg	ggt	gtc	ggt	ata	ttt	tta	ggt

cta cgt gaa ttg gtt cac agt tta ggc atc tct aag cca aca att gta Leu Arg Glu Leu Val His Ser Leu Gly Ile Ser Lys Pro Thr Ile Val 

Gly Leu Phe Ile Gly Val Gly Val Ala Pro Thr Asn Glu Ile Tyr Thr

ttt	agt	tct	aaa	aaa	gga	tta	gat	aaa	gtt	ata	act	gta	caa	aaa	acg	432
Phe	Ser	Ser	Lys	Lys	Gly	Leu	Asp	Lys	Val	Пе	Thr	Val	Gln	Lys	Thr	
	130					135					140					
gta	act	gct	att	aaa	acc	att	gtt	ata	ttg	gac	agc	aaa	gtg	gat	tat	480
Val	Thr	Ala	He	Lys	Thr	I1e	Val	Ile	Leu	Asp	Ser	Lys	Val	Asp	Tyr	
145					150					155					160	
aga	ggt	tat	caa	tcc	atg	gac	aac	ttt	att	aaa	aaa	aac	act	cca	caa	528
Arg	Gly	Tyr	Gln	Ser	Met	Asp	Asn	Phe	lle	Lys	Lys	Asn	Thr	Pro	Gln	
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ggt	ttc	aaa	gga	tca	agt	ttt	aaa	act	gta	gaa	gtt	aac	cgc	aaa	gaa	576
Gly	Phe	Lys	Gly	Ser	Ser	Phe	Lys	Thr	Val	Glu	Val	Asn	Arg	Lys	Glu	
			180					185					190			
caa	gtt	gct	ctt	ata	atg	aac	tct	tcg	ggt	tca	acc	ggt	ttg	cca	aaa	624
Gln	Val	Ala	Leu	Ile	Met	Asn	Ser	Ser	Gly	Ser	Thr	Gly	Leu	Pro	Lys	
		195					200					205				
		caa													_	672
Gly		Gln	Leu	Thr	His	Glu	Asn	He	Val	Thr	Arg	Phe	Ser	His	Ala	
	210					215					220					
		cca														720
Arg	Asp	Pro	He	Tyr		Asn	Gln	Val	Ser		Gly	Thr	Ala	Ile	Leu	
225					230					235					240	

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Thr	Val	Val	Pro	Phe	His	His	Gly	Phe	G1y	Met	Phe	Thr	Thr	Leu	Gly	
				245					250					255		
tat	cta	act	tgt	ggt	ttt	cgt	att	gtc	atg	tta	acg	aaa	ttt	gac	gaa	816
Tyr	Leu	Thr	Cys	Gly	Phe	Arg	Ile	Val	Met	Leu	Thr	Lys	Phe	Asp	Glu	
			260					265					270			
gag	act	ttt	tta	aaa	aca	ctg	caa	gat	tac	aaa	tgt	tca	agc	gtt	att	864
Glu	Thr	Phe	Leu	Lys	Thr	Leu	Gln	Asp	Tyr	Lys	Cys	Ser	Ser	Val	Ile	
		275					280					285				
ctt	gta	ccg	act	ttg	ttt	gca	att	ctt	aat	aga	agt	gaa	tta	ctc	gat	912
Leu	Val	Pro	Thr	Leu	Phe	Ala	lle	Leu	Asn	Arg	Ser	G1 u	Leu	Leu	Asp	
	290					295					300					
	tat															960
Lys	tat Tyr				Asn					Ala					Pro	960
																960
Lys 305	Tyr	Asp	Leu	Ser	Asn 310	Leu	Val	Glu	Ile	Ala 315	Ser	Gly	Gly	Ala	Pro 320	
Lys 305 tta	Tyr	Asp	Leu	Ser	Asn 310 ggt	Leu	Val gct	Glu gtt	Ile gct	Ala 315 aga	Ser	Gly ttt	Gly aat	Ala tta	Pro 320 ccg	960
Lys 305 tta	Tyr	Asp	Leu	Ser att Ile	Asn 310 ggt	Leu	Val gct	Glu gtt	Ile gct Ala	Ala 315 aga	Ser	Gly ttt	Gly aat	Ala tta Leu	Pro 320 ccg	
Lys 305 tta	Tyr	Asp	Leu	Ser	Asn 310 ggt	Leu	Val gct	Glu gtt	Ile gct	Ala 315 aga	Ser	Gly ttt	Gly aat	Ala tta	Pro 320 ccg	
Lys 305 tta Leu	Tyr tct Ser	Asp aaa Lys	Leu gaa Glu	ser att Ile 325	Asn 310 ggt Gly	Leu gaa Glu	Val gct Ala	Glu gtt Val	gct Ala 330	Ala 315 aga Arg	Ser cgt Arg	Gly ttt Phe	Gly aat Asn	Ala tta Leu 335	Pro 320 ccg Pro	1008
Lys 305 tta Leu	tct Ser	Asp aaa Lys	Leu gaa Glu caa	ser att Ile 325	Asn 310 ggt Gly tat	Leu gaa Glu ggt	Val gct Ala	Glu gtt Val aca	gct Ala 330	Ala 315 aga Arg	Ser cgt Arg	ttt Phe	Gly aat Asn	tta Leu 335	Pro 320 ccg Pro	
Lys 305 tta Leu	Tyr tct Ser	Asp aaa Lys	gaa Glu caa Gln	ser att Ile 325	Asn 310 ggt Gly tat	Leu gaa Glu ggt	Val gct Ala	gtt Val aca Thr	gct Ala 330	Ala 315 aga Arg	Ser cgt Arg	ttt Phe	Gly aat Asn gca Ala	tta Leu 335	Pro 320 ccg Pro	1008
Lys 305 tta Leu	tct Ser	Asp aaa Lys	Leu gaa Glu caa	ser att Ile 325	Asn 310 ggt Gly tat	Leu gaa Glu ggt	Val gct Ala	Glu gtt Val aca	gct Ala 330 gaa	Ala 315 aga Arg	Ser cgt Arg	ttt Phe	Gly aat Asn	tta Leu 335	Pro 320 ccg Pro	1008

Ile Thr Pro Glu Gly Asp Asp Lys Pro Gly Ala Ser Gly Lys Val Val
355 360 365

cca	tta	ttt	aaa	gca	aaa	gtt	atc	gat	ctt	gat	act	aaa	aaa	act	ttg	1152
Pro	Leu	Phe	Lys	Ala	Lys	Val	Ile	Asp	Leu	Asp	Thr	Lys	Lys	Thr	Leu	
	370					375					380					

ggc	ccg	aac	aga	cgt	gga	gaa	gtt	tgt	gta	aag	ggt	cct	atg	ctt	atg	1200
Gly	Pro	Asn	Arg	Arg	Gly	Glu	Val	Cys	Val	Lys	Gly	Pro	Met	Leu	Met	
385					390					395					400	

aaa	ggt	tat	gta	gat	aat	cca	gaa	gca	aca	aga	gaa	atc	ata	gat	gaa	1248
Lys	Gly	Tyr	Val	Asp	Asn	Pro	Glu	Ala	Thr	Arg	G1 u	Ile	Ile	Asp	Glu	
				405					410					415		

gaa	ggt	tgg	ttg	cac	aca	gga	gat	att	ggg	tat	tac	gat	gaa	gaa	aaa	]	1296
Glu	Gly	Trp	Leu	His	Thr	Gly	Asp	Ile	Gly	Tyr	Tyr	Asp	G1 u	Glu	Lys		
			420					425					430				

cat	ttc	ttt	atc	gtg	gat	cgt	ttg	aag	tct	tta	atc	aaa	tac	aaa	gga	1	344
His	Phe	Phe	Ile	Val	Asp	Arg	Leu	Lys	Ser	Leu	Ile	Lys	Tyr	Lys	Gly		
		435					440					445					

tat	caa	gta	cca	cct	gct	gaa	tta	gaa	tct	gtt	ctt	ttg	caa	cat	cca	1392
Tyr	Gln	Val	Pro	Pro	Ala	Glu	Leu	Glu	Ser	Val	Leu	Leu	Gln	His	Pro	
	450					455					460					

aat att ttt gat gcc ggc gtt gct ggc gtt cca gat cct ata gct ggt 1440 Asn Ile Phe Asp Ala Gly Val Ala Gly Val Pro Asp Pro Ile Ala Gly

13

465	470	475	480

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gaa aaa gaa gta atg gat tac gtt gct agt caa gtt tca aat gca aaa 1536 Glu Lys Glu Val Met Asp Tyr Val Ala Ser Gln Val Ser Asn Ala Lys 500 505 510

cgt ttg cgt ggt ggt gtc cgt ttt gtg gac gaa gta cct aaa ggt ctc 1584 Arg Leu Arg Gly Gly Val Arg Phe Val Asp Glu Val Pro Lys Gly Leu 515 520 525

act ggt aaa att gac ggt aaa gca att aga gaa ata ctg aag aaa cca 1632 Thr Gly Lys Ile Asp Gly Lys Ala Ile Arg Glu Ile Leu Lys Lys Pro 530 535 540

gtt gct aag atg 1644 Val Ala Lys Met 545

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Met Asp Arg Tyr Ala Lys Leu Gly Ala Ile Ala Phe Thr Asn Ala Leu 35 40 45

Thr Gly Val Asp Tyr Thr Tyr Ala Glu Tyr Leu Glu Lys Ser Cys Cys
50 55 60

Leu Gly Glu Ala Leu Lys Asn Tyr Gly Leu Val Val Asp Gly Arg Ile
65 70 75 80

Ala Leu Cys Ser Glu Asn Cys Glu Glu Phe Phe Ile Pro Val Leu Ala 85 90 95

Gly Leu Phe Ile Gly Val Gly Val Ala Pro Thr Asn Glu Ile Tyr Thr 100 105 110

Leu Arg Glu Leu Val His Ser Leu Gly Ile Ser Lys Pro Thr Ile Val 115 120 125

Phe Ser Ser Lys Lys Gly Leu Asp Lys Val IIe Thr Val Gln Lys Thr
130 135 140

Val Thr Ala Ile Lys Thr Ile Val Ile Leu Asp Ser Lys Val Asp Tyr 145 150 155 160 Arg Gly Tyr Gln Ser Met Asp Asn Phe Ile Lys Lys Asn Thr Pro Gln
165 170 175

Gly Phe Lys Gly Ser Ser Phe Lys Thr Val Glu Val Asn Arg Lys Glu 180 185 190

Gln Val Ala Leu Ile Met Asn Ser Ser Gly Ser Thr Gly Leu Pro Lys 195 200 205

Gly Val Gln Leu Thr His Glu Asn Ile Val Thr Arg Phe Ser His Ala 210 215 220

Arg Asp Pro Ile Tyr Gly Asn Gln Val Ser Pro Gly Thr Ala Ile Leu 225 230 235 240

Thr Val Val Pro Phe His His Gly Phe Gly Met Phe Thr Thr Leu Gly 245 250 255

Tyr Leu Thr Cys Gly Phe Arg Ile Val Met Leu Thr Lys Phe Asp Glu 260 265 270

Glu Thr Phe Leu Lys Thr Leu Gln Asp Tyr Lys Cys Ser Ser Val Ile 275 280 285

Leu Val Pro Thr Leu Phe Ala IIe Leu Asn Arg Ser Glu Leu Leu Asp 290 295 300

Lys Tyr Asp Leu Ser Asn Leu Val Glu Ile Ala Ser Gly Gly Ala Pro

Leu Ser Lys Glu Ile Gly Glu Ala Val Ala Arg Arg Phe Asn Leu Pro 

Gly Val Arg Gln Gly Tyr Gly Leu Thr Glu Thr Thr Ser Ala Ile Ile 

lle Thr Pro Glu Gly Asp Asp Lys Pro Gly Ala Ser Gly Lys Val Val 

Pro Leu Phe Lys Ala Lys Val Ile Asp Leu Asp Thr Lys Lys Thr Leu 

Gly Pro Asn Arg Arg Gly Glu Val Cys Val Lys Gly Pro Met Leu Met 

Lys Gly Tyr Val Asp Asn Pro Glu Ala Thr Arg Glu Ile Ile Asp Glu 

Glu Gly Trp Leu His Thr Gly Asp Ile Gly Tyr Tyr Asp Glu Glu Lys 

His Phe Phe Ile Val Asp Arg Leu Lys Ser Leu Ile Lys Tyr Lys Gly 

Tyr Gln Val Pro Pro Ala Glu Leu Glu Ser Val Leu Leu Gln His Pro 

Asn Ile Phe Asp Ala Gly Val Ala Gly Val Pro Asp Pro Ile Ala Gly
465 470 475 480

Glu Leu Pro Gly Ala Val Val Leu Lys Lys Gly Lys Ser Met Thr
485 490 495

Glu Lys Glu Val Met Asp Tyr Val Ala Ser Gln Val Ser Asn Ala Lys
500 505 510

Arg Leu Arg Gly Gly Val Arg Phe Val Asp Glu Val Pro Lys Gly Leu 515 520 525

Thr Gly Lys Ile Asp Gly Lys Ala Ile Arg Glu Ile Leu Lys Lys Pro 530 535 540

Val Ala Lys Met 545